ABSTRACT OF THE DISCLOSURE

An Arabidopsis thaliana double mutant ssl2 slr having a mutation in at least one base of the SSL2 genomic gene shown in SEQ ID NO: 3, obtained by: treating an Arabidopsis thaliana slr dominant mutant (FERM BP-8385), which has no lateral roots, with a mutagen; preparing plants of the next generation of the mutagen-treated slr dominant mutant; and selecting a plant that basically preserves phenotypes of the slr dominant mutant but has lateral roots from the plants of the next generation.

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